

SEQUENCE LISTING

<110> MEYERS, Rachel A.
WILLIAMSON, Mark

<120> 47169 and 33935, Novel Human Glycosyl Transferases and
Uses Thereof

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<150> US 60/249,939

<151> 2000-11-20

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Arg Asn Pro Phe Leu Ser Lys Val Lys Leu Ile Tyr Tyr Tyr Leu Phe		
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Ala Phe Ile Tyr Gly Leu Val Gly Ser Cys Ser Asp Val Val Met Val		
	245	250 255
Asn Ser Ser Trp Thr Leu Asn His Ile Leu Ser Leu Trp Lys Val Gly		
	260	265 270
Asn Cys Thr Asn Ile Val Tyr Pro Pro Cys Asp Val Gln Thr Phe Leu		
	275	280 285
Asp Ile Pro Leu His Glu Lys Lys Met Thr Pro Gly His Leu Leu Val		
	290	295 300
Ser Val Gly Gln Phe Arg Pro Glu Lys Asn His Pro Leu Gln Ile Arg		

305 310 315 320
 Ala Phe Ala Lys Leu Leu Asn Lys Lys Met Val Glu Ser Pro Pro Ser
 325 330 335
 Leu Lys Leu Val Leu Ile Gly Gly Cys Arg Asn Lys Asp Asp Glu Leu
 340 345 350
 Arg Val Asn Gln Leu Arg Arg Leu Ser Glu Asp Leu Gly Val Gln Glu
 355 360 365
 Tyr Val Glu Phe Lys Ile Asn Ile Pro Phe Asp Glu Leu Lys Asn Tyr
 370 375 380
 Leu Ser Glu Ala Thr Ile Gly Leu His Thr Met Trp Asn Glu His Phe
 385 390 395 400
 Gly Ile Gly Val Val Glu Cys Met Ala Ala Gly Thr Ile Ile Leu Ala
 405 410 415
 His Asn Ser Gly Gly Pro Lys Leu Asp Ile Val Val Pro His Glu Gly
 420 425 430
 Asp Ile Thr Gly Phe Leu Ala Glu Ser Glu Glu Asp Tyr Ala Glu Thr
 435 440 445
 Ile Ala His Ile Leu Ser Met Ser Ala Glu Lys Arg Leu Gln Ile Arg
 450 455 460
 Lys Ser Ala Arg Ala Ser Val Ser Arg Phe Ser Asp Gln Glu Phe Glu
 465 470 475 480
 Val Thr Phe Leu Ser Ser Val Glu Lys Leu Phe Lys
 485 490

<210> 13

<211> 3044

<212> DNA

<213> Homo sapiens

<400> 13

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 aatcaaatgg tgattgcatt ttttcatcca tactgcaatg ctggtggagg aggagaaaga 240
 gttttatggt gtgctttaag agccctgcag aaaaagtatc ctgaagcagt ttatgttggt 300
 tataccggcg atgttaatgt caacgggtcaa cagatactag aaggtgcttt cagaagattt 360

aacatcagat taattcaccc agtgcagttt gtttttttaa ggaaacgcta tcttgtggaa 420
 gattcactgt atcctcactt cactcgtctg ggccaaagtc taggatccat ttttcttggc 480
 tgggaagctc taatgcagtg tgttcctgat gtttacattg attcaatggg atacgctttt 540
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 gccttcatta ccaggaatcc ttttctcagc aaagtaaagc tcatctacta ctatttattt 720
 gcttttattt atggacttgt tggttcttgc agtgatgtag tcatgggtaa ttcttcttgg 780
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 aaaagagact ccaaatcaga aaaagtgtct gtgcatctgt aagcagattc tctgatcagg 3000
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<210> 15
<400> 15
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<210> 18
<400> 18
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<210> 19
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<210> 20
<211> 559
<212> PRT
<213> Homo sapiens

<400> 20
Met Arg Lys Phe Ala Tyr Cys Lys Val Val Leu Ala Thr Ser Leu Ile
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Trp Val Leu Leu Asp Met Phe Leu Leu Leu Tyr Phe Ser Glu Cys Asn
20 25 30

Lys Cys Asp Glu Lys Lys Glu Arg Gly Leu Pro Ala Gly Asp Val Leu
35 40 45

Glu Pro Val Gln Lys Pro His Glu Gly Pro Gly Glu Met Gly Lys Pro
50 55 60

Val Val Ile Pro Lys Glu Asp Gln Glu Lys Met Lys Glu Met Phe Lys

65		70		75		80
Ile Asn Gln Phe Asn Leu Met Ala Ser Glu Met Ile Ala Leu Asn Arg						
	85			90		95
Ser Leu Pro Asp Val Arg Leu Glu Gly Cys Lys Thr Lys Val Tyr Pro						
	100			105		110
Asp Asn Leu Pro Thr Thr Ser Val Val Ile Val Phe His Asn Glu Ala						
	115			120		125
Trp Ser Thr Leu Leu Arg Thr Val His Ser Val Ile Asn Arg Ser Pro						
	130			135		140
Arg His Met Ile Glu Glu Ile Val Leu Val Asp Asp Ala Ser Glu Arg						
	145			150		155
Asp Phe Leu Lys Arg Pro Leu Glu Ser Tyr Val Lys Lys Leu Lys Val						
	165			170		175
Pro Val His Val Ile Arg Met Glu Gln Arg Ser Gly Leu Ile Arg Ala						
	180			185		190
Arg Leu Lys Gly Ala Ala Val Ser Lys Gly Gln Val Ile Thr Phe Leu						
	195			200		205
Asp Ala His Cys Glu Cys Thr Val Gly Trp Leu Glu Pro Leu Leu Ala						
	210			215		220
Arg Ile Lys His Asp Arg Arg Thr Val Val Cys Pro Ile Ile Asp Val						
	225			230		235
Ile Ser Asp Asp Thr Phe Glu Tyr Met Ala Gly Ser Asp Met Thr Tyr						
	245			250		255
Gly Gly Phe Asn Trp Lys Leu Asn Phe Arg Trp Tyr Pro Val Pro Gln						
	260			265		270
Arg Glu Met Asp Arg Arg Lys Gly Asp Arg Thr Leu Pro Val Arg Thr						
	275			280		285
Pro Thr Met Ala Gly Gly Leu Phe Ser Ile Asp Arg Asp Tyr Phe Gln						
	290			295		300
Glu Ile Gly Thr Tyr Asp Ala Gly Met Asp Ile Trp Gly Gly Glu Asn						
	305			310		315
Leu Glu Ile Ser Phe Arg Ile Trp Gln Cys Gly Gly Thr Leu Glu Ile						

325	330	335
Val Thr Cys Ser His Val Gly His Val Phe Arg Lys Ala Thr Pro Tyr		
340	345	350
Thr Phe Pro Gly Gly Thr Gly Gln Ile Ile Asn Lys Asn Asn Arg Arg		
355	360	365
Leu Ala Glu Val Trp Met Asp Glu Phe Lys Asn Phe Phe Tyr Ile Ile		
370	375	380
Ser Pro Gly Val Thr Lys Val Asp Tyr Gly Asp Ile Ser Ser Arg Val		
385	390	395
Gly Leu Arg His Lys Leu Gln Cys Lys Pro Phe Ser Trp Tyr Leu Glu		
405	410	415
Asn Ile Tyr Pro Asp Ser Gln Ile Pro Arg His Tyr Phe Ser Leu Gly		
420	425	430
Glu Ile Arg Lys Glu Glu Thr Asn Gln Cys Leu Asp Asn Met Ala Arg		
435	440	445
Lys Glu Asn Glu Lys Val Gly Ile Phe Asn Cys His Gly Met Gly Gly		
450	455	460
Asn Gln Val Phe Ser Tyr Thr Ala Asn Lys Glu Ile Arg Thr Asp Asp		
465	470	475
Leu Cys Leu Asp Val Ser Lys Leu Asn Gly Pro Val Thr Met Leu Lys		
485	490	495
Cys His His Leu Lys Gly Asn Gln Leu Trp Glu Tyr Asp Pro Val Lys		
500	505	510
Leu Thr Leu Gln His Val Asn Ser Asn Gln Cys Leu Asp Lys Ala Thr		
515	520	525
Glu Glu Asp Ser Gln Val Pro Ser Ile Arg Asp Cys Asn Gly Ser Arg		
530	535	540
Ser Gln Gln Trp Leu Leu Arg Asn Val Thr Leu Pro Glu Ile Phe		
545	550	555

<210> 21
 <211> 559
 <212> PRT

<213> Rattus sp.

<400> 21

Met Arg Lys Phe Ala Tyr Cys Lys Val Val Leu Ala Thr Ser Leu Val
1 5 10 15

Trp Val Leu Leu Asp Met Phe Leu Leu Leu Tyr Phe Ser Glu Cys Asn
20 25 30

Lys Cys Glu Glu Lys Lys Glu Arg Gly Leu Pro Ala Gly Asp Val Leu
35 40 45

Glu Leu Val Gln Lys Pro His Glu Gly Pro Gly Glu Met Gly Lys Pro
50 55 60

Val Val Ile Pro Lys Glu Asp Gln Glu Lys Met Lys Glu Met Phe Lys
65 70 75 80

Ile Asn Gln Phe Asn Leu Met Ala Ser Glu Met Ile Ala Phe Asn Arg
85 90 95

Ser Leu Pro Asp Val Arg Leu Glu Gly Cys Lys Thr Lys Val Tyr Pro
100 105 110

Asp Ser Leu Pro Thr Thr Ser Val Val Ile Val Phe His Asn Glu Ala
115 120 125

Trp Ser Thr Leu Leu Arg Thr Val His Ser Val Ile Asn Arg Ser Pro
130 135 140

Arg His Met Ile Glu Glu Ile Val Leu Val Asp Asp Ala Ser Glu Arg
145 150 155 160

Asp Phe Leu Lys Arg Pro Leu Glu Ser Tyr Val Lys Lys Leu Lys Val
165 170 175

Pro Val His Val Ile Arg Met Glu Gln Arg Ser Gly Leu Ile Arg Ala
180 185 190

Arg Leu Lys Gly Ala Ala Val Ser Lys Gly Gln Val Ile Thr Phe Leu
195 200 205

Asp Ala His Cys Glu Cys Thr Val Gly Trp Leu Glu Pro Leu Leu Ala
210 215 220

Arg Ile Lys His Asp Arg Arg Thr Val Val Cys Pro Ile Ile Asp Val
225 230 235 240

Ile Ser Asp Asp Thr Phe Glu Tyr Met Ala Gly Ser Asp Met Thr Tyr
 245 250 255

Gly Gly Phe Asn Trp Lys Leu Asn Phe Arg Trp Tyr Pro Val Pro Gln
 260 265 270

Arg Glu Met Asp Arg Arg Lys Gly Asp Arg Thr Leu Pro Val Arg Thr
 275 280 285

Pro Thr Met Ala Gly Gly Leu Phe Ser Ile Asp Arg Asp Tyr Phe Gln
 290 295 300

Glu Ile Gly Thr Tyr Asp Ala Gly Met Asp Ile Trp Gly Gly Glu Asn
 305 310 315 320

Leu Glu Ile Ser Phe Arg Ile Trp Gln Cys Gly Gly Thr Leu Glu Ile
 325 330 335

Val Thr Cys Ser His Val Gly His Val Phe Arg Lys Ala Thr Pro Tyr
 340 345 350

Thr Phe Pro Gly Gly Thr Gly Gln Ile Ile Asn Lys Asn Asn Arg Arg
 355 360 365

Leu Ala Glu Val Trp Met Asp Glu Phe Lys Asn Phe Phe Tyr Ile Ile
 370 375 380

Ser Pro Gly Val Thr Lys Val Asp Tyr Gly Asp Ile Ser Ser Arg Val
 385 390 395 400

Gly Leu Arg His Lys Leu Gln Cys Lys Pro Phe Ser Trp Tyr Leu Glu
 405 410 415

Asn Ile Tyr Pro Asp Ser Gln Ile Pro Arg His Tyr Phe Ser Leu Gly
 420 425 430

Glu Ile Arg Asn Val Glu Thr Asn Gln Cys Leu Asp Asn Met Ala Arg
 435 440 445

Lys Glu Asn Glu Lys Val Gly Ile Phe Asn Cys His Gly Met Gly Gly
 450 455 460

Asn Gln Val Phe Ser Tyr Thr Ala Asn Lys Glu Ile Arg Thr Asp Asp
 465 470 475 480

Leu Cys Leu Asp Val Ser Lys Leu Asn Gly Pro Val Thr Met Leu Lys
 485 490 495

Cys His His Leu Lys Gly Asn Gln Leu Trp Glu Tyr Asp Pro Val Lys
 500 505 510

Leu Thr Leu Gln His Val Asn Ser Asn Gln Cys Leu Asp Lys Ala Thr
 515 520 525

Glu Glu Asp Ser Gln Val Pro Ser Ile Arg Asp Cys Thr Gly Ser Arg
 530 535 540

Ser Gln Gln Trp Leu Leu Arg Asn Val Thr Leu Pro Glu Ile Phe
 545 550 555

<210> 22

<211> 559

<212> PRT

<213> Mus sp.

<400> 22

Met Arg Lys Phe Ala Tyr Cys Lys Val Val Leu Ala Thr Ser Leu Val
 1 5 10 15

Trp Val Leu Leu Asp Met Phe Leu Leu Leu Tyr Phe Ser Glu Cys Asn
 20 25 30

Lys Cys Glu Glu Lys Gln Glu Arg Gly Leu Pro Ala Gly Asp Val Leu
 35 40 45

Glu Leu Val Gln Lys Pro His Glu Gly Pro Gly Glu Met Gly Lys Pro
 50 55 60

Val Val Ile Pro Lys Glu Asp Gln Glu Lys Met Lys Glu Met Phe Lys
 65 70 75 80

Ile Asn Gln Phe Asn Leu Met Ala Ser Glu Met Ile Ala Leu Asn Arg
 85 90 95

Ser Leu Pro Asp Val Arg Leu Glu Gly Cys Lys Thr Lys Val Tyr Pro
 100 105 110

Asp Asn Leu Pro Thr Thr Ser Val Val Ile Val Phe His Asn Glu Ala
 115 120 125

Trp Ser Thr Leu Leu Arg Thr Val His Ser Val Ile Asn Arg Ser Pro
 130 135 140

Arg His Met Ile Glu Glu Ile Val Leu Val Asp Asp Ala Ser Glu Arg
 145 150 155 160

Asp Phe Leu Lys Arg Pro Leu Glu Ser Tyr Val Lys Lys Leu Lys Val
 165 170 175

Pro Val His Val Ile Arg Met Glu Gln Arg Ser Gly Leu Ile Arg Ala
 180 185 190

Arg Leu Lys Gly Ala Ala Val Ser Arg Gly Gln Val Ile Thr Phe Leu
 195 200 205

Asp Ala His Cys Glu Cys Thr Ala Gly Trp Leu Glu Pro Leu Leu Ala
 210 215 220

Arg Ile Lys His Asp Arg Arg Thr Val Val Cys Pro Ile Ile Asp Val
 225 230 235 240

Ile Ser Asp Asp Thr Phe Glu Tyr Met Ala Gly Ser Asp Met Thr Tyr
 245 250 255

Gly Gly Phe Asn Trp Lys Leu Asn Phe Arg Trp Tyr Pro Val Pro Gln
 260 265 270

Arg Glu Met Asp Arg Arg Lys Gly Asp Arg Thr Leu Pro Val Arg Thr
 275 280 285

Pro Thr Met Ala Gly Gly Leu Phe Ser Ile Asp Arg Asp Tyr Phe Gln
 290 295 300

Glu Ile Gly Thr Tyr Asp Ala Gly Met Asp Ile Trp Gly Gly Glu Asn
 305 310 315 320

Leu Glu Ile Ser Phe Arg Ile Trp Gln Cys Gly Gly Thr Leu Glu Ile
 325 330 335

Val Thr Cys Ser His Val Gly His Val Phe Arg Lys Ala Thr Pro Tyr
 340 345 350

Thr Phe Pro Gly Gly Thr Gly Gln Ile Ile Asn Lys Asn Asn Arg Arg
 355 360 365

Leu Ala Glu Val Trp Met Asp Glu Phe Lys Asn Phe Phe Tyr Ile Ile
 370 375 380

Ser Pro Gly Val Thr Lys Val Asp Tyr Gly Asp Ile Ser Ser Arg Leu
 385 390 395 400

Gly Leu Arg Arg Lys Leu Gln Cys Lys Pro Phe Ser Trp Tyr Leu Glu
 405 410 415

Asn Ile Tyr Pro Asp Ser Gln Ile Pro Arg His Tyr Phe Ser Leu Gly
420 425 430

Glu Ile Arg Asn Val Glu Thr Asn Gln Cys Leu Asp Asn Met Ala Arg
435 440 445

Lys Glu Asn Glu Lys Val Gly Ile Phe Asn Cys His Gly Met Gly Gly
450 455 460

Asn Gln Val Phe Ser Tyr Thr Ala Asn Lys Glu Ile Arg Thr Asp Asp
465 470 475 480

Leu Cys Leu Asp Val Ser Lys Leu Asn Gly Pro Val Thr Met Leu Lys
485 490 495

Cys His His Leu Lys Gly Asn Gln Leu Trp Glu Tyr Asp Pro Val Lys
500 505 510

Leu Thr Leu Gln His Val Asn Ser Asn Gln Cys Leu Asp Lys Ala Thr
515 520 525

Glu Glu Asp Ser Gln Val Pro Ser Ile Arg Asp Cys Thr Gly Ser Arg
530 535 540

Ser Gln Gln Trp Leu Leu Arg Asn Val Thr Leu Pro Glu Ile Phe
545 550 555

<210> 23

<211> 559

<212> PRT

<213> Bos sp.

<400> 23

Met Arg Lys Phe Ala Tyr Cys Lys Val Val Leu Ala Thr Ser Leu Ile
1 5 10 15

Trp Val Leu Leu Asp Met Phe Leu Leu Leu Tyr Phe Ser Glu Cys Asn
20 25 30

Lys Cys Asp Glu Lys Lys Glu Arg Gly Leu Pro Ala Gly Asp Val Leu
35 40 45

Glu Pro Val Gln Lys Pro His Glu Gly Pro Gly Glu Met Gly Lys Pro
50 55 60

Val Val Ile Pro Lys Glu Asp Gln Glu Lys Met Lys Glu Met Phe Lys

65		70		75		80
Ile Asn Gln Phe Asn Leu Met Ala Ser Glu Met Ile Ala Leu Asn Arg						
	85		90		95	
Ser Leu Pro Asp Val Arg Leu Glu Gly Cys Lys Thr Lys Val Tyr Pro						
	100		105		110	
Asp Asn Leu Pro Thr Thr Ser Val Val Ile Val Phe His Asn Glu Ala						
	115		120		125	
Trp Ser Thr Leu Leu Arg Thr Val His Ser Val Ile Asn Arg Ser Pro						
	130		135		140	
Arg His Met Leu Glu Glu Ile Val Leu Val Asp Asp Ala Ser Glu Arg						
	145		150		155	160
Asp Phe Leu Lys Arg Pro Leu Glu Ser Tyr Val Lys Lys Leu Lys Val						
		165		170		175
Pro Val His Val Ile Arg Met Glu Gln Arg Ser Gly Leu Ile Arg Ala						
	180		185		190	
Arg Leu Lys Gly Ala Ala Val Ser Lys Gly Gln Val Ile Thr Phe Leu						
	195		200		205	
Asp Ala His Cys Glu Cys Thr Val Gly Trp Leu Glu Pro Leu Leu Ala						
	210		215		220	
Arg Ile Lys His Asp Arg Lys Thr Val Val Cys Pro Ile Ile Asp Val						
	225		230		235	240
Ile Ser Asp Asp Thr Phe Glu Tyr Met Ala Gly Ser Asp Met Thr Tyr						
	245		250		255	
Gly Gly Phe Asn Trp Lys Leu Asn Phe Arg Trp Tyr Pro Val Pro Gln						
	260		265		270	
Arg Glu Met Asp Arg Arg Lys Gly Asp Arg Thr Leu Pro Val Arg Thr						
	275		280		285	
Pro Thr Met Ala Gly Gly Leu Phe Ser Ile Asp Arg Asp Tyr Phe Gln						
	290		295		300	
Glu Ile Gly Thr Tyr Asp Ala Gly Met Asp Ile Trp Gly Gly Glu Asn						
	305		310		315	320
Leu Glu Ile Ser Phe Arg Ile Trp Gln Cys Gly Gly Thr Leu Glu Ile						

Val Thr Cys Ser His Val Gly His Val Phe Arg Lys Ala Thr Pro Tyr	325	330	335
340	345	350	
Thr Phe Pro Gly Gly Thr Gly Gln Ile Ile Asn Lys Asn Asn Arg Arg	355	360	365
Leu Ala Glu Val Trp Met Asp Glu Phe Lys Asn Phe Phe Tyr Ile Ile	370	375	380
Ser Pro Gly Val Thr Lys Val Asp Tyr Gly Asp Ile Ser Ser Arg Leu	385	390	395
400			
Gly Leu Arg His Lys Leu Gln Cys Arg Pro Phe Ser Trp Tyr Leu Glu	405	410	415
Asn Ile Tyr Pro Asp Ser Gln Ile Pro Arg His Tyr Phe Ser Leu Gly	420	425	430
Glu Ile Arg Asn Val Glu Thr Asn Gln Cys Leu Asp Asn Met Ala Arg	435	440	445
Lys Glu Asn Glu Lys Val Gly Ile Phe Asn Cys His Gly Met Gly Gly	450	455	460
Asn Gln Val Phe Ser Tyr Thr Ala Asn Lys Glu Ile Arg Thr Asp Asp	465	470	475
480			
Leu Cys Leu Asp Val Ser Lys Leu Asn Gly Pro Val Thr Met Leu Lys	485	490	495
Cys His His Leu Lys Gly Asn Gln Leu Trp Glu Tyr Asp Pro Val Lys	500	505	510
Leu Thr Leu Gln His Val Asn Ser Asn Gln Cys Leu Asp Lys Ala Thr	515	520	525
Asp Glu Asp Ser Gln Val Pro Ser Ile Arg Asp Cys Ser Gly Ser Arg	530	535	540
Ser Gln Gln Trp Leu Leu Arg Asn Val Thr Leu Pro Glu Ile Phe	545	550	555

<210> 24
 <211> 559
 <212> PRT

<213> Sus sp.

<400> 24

Met Arg Lys Phe Ala Tyr Cys Lys Val Val Leu Ala Thr Ser Leu Ile
1 5 10 15

Trp Val Leu Leu Asp Met Phe Leu Leu Leu Tyr Phe Ser Glu Cys Asn
20 25 30

Lys Cys Asp Glu Lys Lys Glu Arg Gly Leu Pro Ala Gly Asp Val Leu
35 40 45

Glu Pro Val Gln Lys Pro His Glu Gly Pro Gly Glu Met Gly Lys Pro
50 55 60

Val Val Ile Pro Lys Glu Asp Gln Asp Lys Met Lys Glu Met Phe Lys
65 70 75 80

Ile Asn Gln Phe Asn Leu Met Ala Ser Glu Met Ile Ala Leu Asn Arg
85 90 95

Ser Leu Pro Asp Val Arg Leu Glu Gly Cys Lys Thr Lys Val Tyr Pro
100 105 110

Asp Asn Leu Pro Thr Thr Ser Val Val Ile Val Phe His Asn Glu Ala
115 120 125

Trp Ser Thr Leu Leu Arg Thr Val His Ser Val Ile Asn Arg Ser Pro
130 135 140

Arg His Met Leu Glu Glu Ile Val Leu Val Asp Asp Ala Ser Glu Arg
145 150 155 160

Asp Phe Leu Lys Arg Pro Leu Glu Ser Tyr Val Lys Lys Leu Lys Val
165 170 175

Pro Val His Val Ile Arg Met Glu Gln Arg Ser Gly Leu Ile Arg Ala
180 185 190

Arg Leu Lys Gly Ala Ala Val Ser Lys Gly Gln Val Ile Thr Phe Leu
195 200 205

Asp Ala His Cys Glu Cys Thr Val Gly Trp Leu Glu Pro Leu Leu Ala
210 215 220

Arg Ile Lys His Asp Arg Lys Thr Val Val Cys Pro Ile Ile Asp Val
225 230 235 240

Ile Ser Asp Asp Thr Phe Glu Tyr Met Ala Gly Ser Asp Met Thr Tyr
 245 250 255
 Gly Gly Phe Asn Trp Lys Leu Asn Phe Arg Trp Tyr Pro Val Pro Gln
 260 265 270
 Arg Glu Met Asp Arg Arg Lys Gly Asp Arg Thr Leu Pro Val Arg Thr
 275 280 285
 Pro Thr Met Ala Gly Gly Leu Phe Ser Ile Asp Arg Asp Tyr Phe Gln
 290 295 300
 Glu Ile Gly Thr Tyr Asp Ala Gly Met Asp Ile Trp Gly Gly Glu Asn
 305 310 315 320
 Leu Glu Ile Ser Phe Arg Ile Trp Gln Cys Gly Gly Thr Leu Glu Ile
 325 330 335
 Val Thr Cys Ser His Val Gly His Val Phe Arg Lys Ala Thr Pro Tyr
 340 345 350
 Thr Phe Pro Gly Gly Thr Gly Gln Ile Ile Asn Lys Asn Asn Arg Arg
 355 360 365
 Leu Ala Glu Val Trp Met Asp Glu Phe Lys Thr Phe Phe Tyr Ile Ile
 370 375 380
 Ser Pro Gly Val Thr Lys Val Asp Tyr Gly Asp Ile Ser Ser Arg Leu
 385 390 395 400
 Gly Leu Arg His Lys Leu Gln Cys Arg Pro Phe Ser Trp Tyr Leu Glu
 405 410 415
 Asn Ile Tyr Pro Asp Ser Gln Ile Pro Arg His Tyr Ser Ser Leu Gly
 420 425 430
 Glu Ile Arg Asn Val Glu Thr Asn Gln Cys Leu Asp Asn Met Ala Arg
 435 440 445
 Lys Glu Asn Glu Lys Val Gly Ile Phe Asn Cys His Gly Met Gly Gly
 450 455 460
 Asn Gln Val Phe Ser Tyr Thr Ala Asn Lys Glu Ile Arg Thr Asp Asp
 465 470 475 480
 Leu Cys Leu Asp Val Ser Lys Leu Asn Gly Pro Val Thr Met Leu Lys
 485 490 495

Cys His His Leu Lys Gly Asn Gln Leu Trp Glu Tyr Asp Pro Val Lys
500 505 510

Leu Thr Leu Gln His Val Asn Ser Asn Gln Cys Leu Asp Lys Ala Thr
515 520 525

Glu Glu Asp Ser Gln Val Pro Ser Ile Arg Asp Cys Ser Gly Ser Arg
530 535 540

Ser Gln Gln Trp Leu Leu Arg Asn Val Thr Leu Pro Glu Ile Phe
545 550 555

<210> 25

<211> 612

<212> PRT

<213> Caenorhabditis elegans

<400> 25

Met Leu Ser Val Gly Gly Gly Arg Ser Ala Val Cys Arg Ala Val Ile
1 5 10 15

Ala Thr Ser Ile Val Trp Leu Leu Ile Asp Val Val Ile Leu Phe Tyr
20 25 30

Tyr Leu Asp Pro Ser Thr Ser Gln Gln Gln Pro Phe Pro Glu Asp Asn
35 40 45

Arg Ile Leu Asn Arg Ala Arg Arg Ile Glu Pro Leu Pro Pro Ala Ala
50 55 60

Gln His Asp Ser Asp Pro Asp Ala His Pro Ile Gln Pro Glu Lys Gln
65 70 75 80

Glu Lys Gln Val Tyr Pro Val Asp Lys Glu Thr Ala Asn Gln Leu Arg
85 90 95

Lys Leu Met Glu Thr Gln Ala Phe Gly Pro Gly Tyr His Gly Gln Gly
100 105 110

Gly Thr Gly Val Thr Val Pro Glu Asp Lys Lys Thr Ile Lys Glu Lys
115 120 125

Arg Phe Leu Glu Asn Gln Phe Asn Val Val Ala Ser Glu Met Ile Ser
130 135 140

Val Asn Arg Thr Leu Pro Asp Tyr Arg Ser Asp Ala Cys Arg Thr Ser
145 150 155 160

Gly Asn Asn Leu Lys Thr Ala Gly Met Pro Lys Thr Ser Ile Ile Ile
 165 170 175
 Val Phe His Asn Glu Ala Trp Thr Thr Leu Leu Arg Thr Leu His Ser
 180 185 190
 Val Ile Asn Arg Ser Pro Arg His Leu Leu Glu Glu Ile Ile Leu Val
 195 200 205
 Asp Asp Lys Ser Asp Arg Asp Tyr Leu Val Lys Pro Leu Asp Ser Tyr
 210 215 220
 Ile Lys Met Phe Pro Ile Pro Ile His Leu Val His Leu Glu Asn Arg
 225 230 235 240
 Ser Gly Leu Ile Arg Ala Arg Leu Thr Gly Ser Glu Met Ala Lys Gly
 245 250 255
 Lys Ile Leu Leu Phe Leu Asp Ala His Val Glu Val Thr Asp Gly Trp
 260 265 270
 Leu Glu Pro Leu Val Ser Arg Val Ala Glu Asp Arg Lys Arg Val Val
 275 280 285
 Ala Pro Ile Ile Asp Val Ile Ser Asp Asp Thr Phe Glu Tyr Val Thr
 290 295 300
 Ala Ser Glu Thr Thr Trp Gly Gly Phe Asn Trp His Leu Asn Phe Arg
 305 310 315 320
 Trp Tyr Ala Val Pro Lys Arg Glu Leu Asn Arg Arg Gly Ser Asp Arg
 325 330 335
 Ser Met Pro Ile Gln Thr Pro Thr Ile Ala Gly Gly Leu Phe Ala Ile
 340 345 350
 Asp Lys Gln Phe Phe Tyr Asp Ile Gly Ser Tyr Asp Glu Gly Met Gln
 355 360 365
 Val Trp Gly Gly Glu Asn Leu Glu Ile Ser Phe Arg Val Trp Met Cys
 370 375 380
 Gly Gly Ser Leu Glu Ile His Pro Cys Ser Arg Val Gly His Val Phe
 385 390 395 400
 Arg Lys Gln Thr Pro Tyr Thr Phe Pro Gly Gly Thr Ala Lys Val Ile
 405 410 415

His His Asn Ala Ala Arg Thr Ala Glu Val Trp Met Asp Glu Tyr Lys
 420 425 430

Ala Phe Phe Tyr Lys Met Val Pro Ala Ala Arg Asn Val Glu Ala Gly
 435 440 445

Asp Val Ser Glu Arg Lys Lys Leu Arg Glu Thr Leu Gln Cys Lys Ser
 450 455 460

Phe Lys Trp Tyr Leu Glu Asn Ile Tyr Pro Glu Ala Pro Leu Pro Ala
 465 470 475 480

Asp Phe Arg Ser Leu Gly Ala Ile Val Asn Arg Phe Thr Glu Lys Cys
 485 490 495

Val Asp Thr Asn Gly Lys Lys Asp Gly Gln Ala Pro Gly Ile Gln Ala
 500 505 510

Cys His Gly Ala Gly Gly Asn Gln Ala Trp Ser Leu Thr Gly Lys Gly
 515 520 525

Glu Ile Arg Ser Asp Asp Leu Cys Leu Ser Ser Gly His Val Tyr Gln
 530 535 540

Ile Gly Ser Glu Leu Lys Leu Glu Arg Cys Ser Val Ser Lys Ile Asn
 545 550 555 560

Val Lys His Val Phe Val Phe Asp Asp Gln Ala Gly Thr Leu Leu His
 565 570 575

Lys Lys Thr Gly Lys Cys Val Thr Gly Ala Asp Gln Arg Val Thr Leu
 580 585 590

Asp Glu Cys Gly Leu Gly Arg Lys Asp Gln Met Trp Gln Leu Glu Gly
 595 600 605

Tyr Gln Ser Pro
 610

<210> 26
 <211> 589
 <212> PRT
 <213> Caenorhabditis elegans

<400> 26
 Met Leu Pro Arg Met Leu Lys Met Lys Thr Val Gly Thr Val Leu Ala

1	5	10	15
Val Ile Trp Leu Phe Gly Leu Ala Phe Ile Tyr Val Gln Ser Thr Ser	20	25	30
Ser Ser Leu Arg Pro Pro Gly Arg His Pro Pro Pro Leu Pro Gln Leu	35	40	45
Asp Pro Leu Ile Pro Gln Asn Pro Pro Gln Asn Asp Glu Ile Arg Pro	50	55	60
Lys Lys Ser Ala Pro Pro Ile Pro Thr Ile Asn Leu Ala Glu Asp Thr	65	70	75
Thr Ile His Glu Arg Thr Glu Lys Asp Val Thr Trp Lys Thr Phe Asp	85	90	95
Val Glu Lys Phe Leu Asn Lys Gly Lys Trp His Gln Gly Glu Asp Lys	100	105	110
Tyr Lys Ala Asn Ser Phe Asn Gln Glu Ala Ser Asp Ala Leu Asn Pro	115	120	125
Thr Arg Lys Ile Pro Asp Ser Arg Glu Pro Gln Cys Arg Asp Val Asp	130	135	140
Tyr Ser Lys Val Gly Met Gln Pro Thr Thr Val Ile Ile Thr Tyr His	145	150	155
Asn Glu Ala Arg Ser Ser Leu Leu Arg Thr Val Phe Ser Val Phe Asn	165	170	175
Gln Ser Pro Glu Glu Leu Leu Leu Glu Ile Val Leu Val Asp Asp Asn	180	185	190
Ser Gln Asp Val Glu Ile Gly Lys Glu Leu Ala Gln Ile Gln Arg Ile	195	200	205
Thr Val Leu Arg Asn Asn Gln Arg Glu Gly Leu Ile Arg Ser Arg Val	210	215	220
Lys Gly Ala Gln Val Ala Arg Ala Pro Val Leu Thr Phe Leu Asp Ser	225	230	235
His Ile Glu Cys Asn Gln Lys Trp Leu Glu Pro Leu Leu Ala Arg Ile	245	250	255
Ala Glu Asn Pro Lys Ala Val Val Ala Pro Ile Ile Asp Val Ile Asn			

260	265	270
Val Asp Asn Phe Asn Tyr Val Gly Ala Ser Ala Asp Leu Arg Gly Gly		
275	280	285
Phe Asp Trp Thr Leu Val Phe Arg Trp Glu Phe Met Asn Glu Gln Leu		
290	295	300
Arg Lys Glu Arg His Ala His Pro Thr Ala Pro Ile Arg Ser Pro Thr		
305	310	315
Met Ala Gly Gly Leu Phe Ala Ile Ser Lys Glu Trp Phe Asn Glu Leu		
	325	330
Gly Thr Tyr Asp Leu Asp Met Glu Val Trp Gly Gly Glu Asn Leu Glu		
	340	345
Met Ser Phe Arg Val Trp Gln Cys Gly Gly Ser Leu Glu Ile Met Pro		
	355	360
Cys Ser Arg Val Gly His Val Phe Arg Lys Lys His Pro Tyr Thr Phe		
	370	375
Pro Gly Gly Ser Gly Asn Val Phe Gln Lys Asn Thr Arg Arg Ala Ala		
	385	390
Glu Val Trp Met Asp Glu Tyr Lys Ala Ile Tyr Leu Lys Asn Val Pro		
	405	410
Ser Ala Arg Phe Val Asn Phe Gly Asp Ile Thr Asp Arg Leu Ala Ile		
	420	425
Arg Asp Arg Leu Gln Cys Lys Ser Phe Lys Trp Tyr Leu Glu Asn Val		
	435	440
Tyr Pro Gln Leu Glu Ile Pro Arg Lys Thr Pro Gly Lys Ser Phe Gln		
	450	455
Met Lys Ile Gly Asn Leu Cys Leu Asp Ser Met Ala Arg Lys Glu Ser		
	465	470
Glu Ala Pro Gly Leu Phe Gly Cys His Gly Thr Gly Gly Asn Gln Glu		
	485	490
Trp Val Phe Asp Gln Leu Thr Lys Thr Phe Lys Asn Ala Ile Ser Gln		
	500	505
Leu Cys Leu Asp Phe Ser Ser Asn Thr Glu Asn Lys Thr Val Thr Met		

515 520 525
 Val Lys Cys Glu Asn Leu Arg Pro Asp Thr Met Val Val Glu Lys Asn
 530 535 540
 Gly Trp Leu Thr Gln Gly Gly Lys Cys Leu Thr Val Asn Gln Gly Ser
 545 550 555 560
 Gly Gly Asp Trp Leu Ile Tyr Gly Ala His Cys Glu Leu Asn Asn Gly
 565 570 575
 Ala Gln Arg Trp Ile Phe Glu Lys Leu Asp Thr Tyr Glu
 580 585

<210> 27
 <211> 626
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 27
 Met Ile Ile Phe Lys Lys Lys Ala Ile Leu Lys Val Leu Leu Leu Val
 1 5 10 15
 Pro Val Phe Trp Ile Cys Ser Leu Ile Phe Phe Ala Ala Thr Ser Asn
 20 25 30
 Asp Ser Ser Gln Ile Gly Ser Asn Asn Asp Leu Ala Asn Lys Ile Ala
 35 40 45
 Glu Ala Asn Phe His Pro Lys Ala Ala Lys Gln Asp Val Ile Gln Gly
 50 55 60
 Phe Gly Pro Pro Ile Glu Pro Glu Pro Val Val Glu Asn Asn Lys Val
 65 70 75 80
 Glu Glu Glu Glu Gln Pro Gly Gly Asn Leu Ala Lys Pro Lys Phe Met
 85 90 95
 Val Asp Pro Asn Asp Pro Ile Tyr Lys Lys Gly Asp Ala Ala Gln Ala
 100 105 110
 Gly Glu Leu Gly Lys Ala Val Val Val Asp Lys Thr Lys Leu Ser Thr
 115 120 125
 Glu Glu Lys Ala Lys Tyr Asp Lys Gly Met Leu Asn Asn Ala Phe Asn
 130 135 140

Gln Tyr Ala Ser Asp Met Ile Ser Val His Arg Thr Leu Pro Thr Asn
145 150 155 160

Ile Asp Ala Glu Cys Lys Thr Glu Lys Tyr Asn Glu Asn Leu Pro Arg
165 170 175

Thr Ser Val Ile Ile Cys Phe His Asn Glu Ala Trp Ser Val Leu Leu
180 185 190

Arg Thr Val His Ser Val Leu Glu Arg Thr Pro Asp His Leu Leu Glu
195 200 205

Glu Val Val Leu Val Asp Asp Phe Ser Asp Met Asp His Thr Lys Arg
210 215 220

Pro Leu Glu Glu Tyr Met Ser Gln Phe Gly Gly Lys Val Lys Ile Leu
225 230 235 240

Arg Met Glu Lys Arg Glu Gly Leu Ile Arg Ala Arg Leu Arg Gly Ala
245 250 255

Ala Val Ala Thr Gly Glu Val Leu Thr Tyr Leu Asp Ser His Cys Glu
260 265 270

Cys Met Glu Gly Trp Met Glu Pro Leu Leu Asp Arg Ile Lys Arg Asp
275 280 285

Pro Thr Thr Val Val Cys Pro Val Ile Asp Val Ile Asp Asp Asn Thr
290 295 300

Phe Glu Tyr His His Ser Lys Ala Tyr Phe Thr Ser Val Gly Gly Phe
305 310 315 320

Asp Trp Gly Leu Gln Phe Asn Trp His Ser Ile Pro Glu Arg Asp Arg
325 330 335

Lys Asn Arg Thr Arg Pro Ile Asp Pro Val Arg Ser Pro Thr Met Ala
340 345 350

Gly Gly Leu Phe Ser Ile Asp Lys Glu Tyr Phe Glu Lys Leu Gly Thr
355 360 365

Tyr Asp Pro Gly Phe Asp Ile Trp Gly Gly Glu Asn Leu Glu Leu Ser
370 375 380

Phe Lys Ile Trp Met Cys Gly Gly Thr Leu Glu Ile Val Pro Cys Ser
385 390 395 400

His Val Gly His Val Phe Arg Lys Arg Ser Pro Tyr Lys Trp Arg Thr
 405 410 415

Gly Val Asn Val Leu Lys Arg Asn Ser Ile Arg Leu Ala Glu Val Trp
 420 425 430

Leu Asp Asp Tyr Lys Thr Tyr Tyr Tyr Glu Arg Ile Asn Asn Gln Leu
 435 440 445

Gly Asp Phe Gly Asp Ile Ser Ser Arg Lys Lys Leu Arg Glu Asp Leu
 450 455 460

Gly Cys Lys Ser Phe Lys Trp Tyr Leu Asp Asn Ile Tyr Pro Glu Leu
 465 470 475 480

Phe Val Pro Gly Glu Ser Val Ala Lys Gly Glu Val Arg Asn Ser Ala
 485 490 495

Val Gln Pro Ala Arg Cys Leu Asp Cys Met Val Gly Arg His Glu Lys
 500 505 510

Asn Arg Pro Val Gly Thr Tyr Gln Cys His Gly Gln Gly Gly Asn Gln
 515 520 525

Tyr Trp Met Leu Ser Lys Asp Gly Glu Ile Arg Arg Asp Glu Ser Cys
 530 535 540

Val Asp Tyr Ala Gly Ser Asp Val Met Val Phe Pro Cys His Gly Met
 545 550 555 560

Lys Gly Asn Gln Glu Trp Arg Tyr Asn His Asp Thr Gly Arg Leu Gln
 565 570 575

His Ala Val Ser Gln Lys Cys Leu Gly Met Thr Lys Asp Gly Ala Lys
 580 585 590

Leu Glu Met Val Ala Cys Gln Tyr Asp Asp Pro Tyr Gln His Trp Lys
 595 600 605

Phe Lys Glu Tyr Asn Glu Ala Lys Ala Ile Glu His Gly Ala Lys Pro
 610 615 620

Pro Ser
 625

<210> 28

<211> 618

<212> PRT

<213> Caenorhabditis elegans

<400> 28

Met Ile Ala Ser Leu Ile Arg Ser Arg Arg Arg Ser Arg Arg Cys Val
1 5 10 15

Val Tyr Ser Val Phe Leu Phe Gly Phe Leu Ala Leu Trp Gly Ser Phe
20 25 30

Ala Leu Ala Leu Val Phe Leu Ser Asp Met Tyr Ile Gly Glu Asp Gln
35 40 45

Ile Ser Thr Gln Lys Ala Ile Lys Pro Ile Ala Arg Ser Asn Tyr His
50 55 60

Val Val Val Gly His Tyr Asn Gly Asn Leu Pro Glu Asp Lys Lys Arg
65 70 75 80

Asn Leu Thr Ser Glu Glu Leu Asn Ala Asn Leu Tyr Ala Pro His Asp
85 90 95

Asp Trp Gly Glu Gly Gly Ala Gly Val Ser His Leu Thr Pro Glu Gln
100 105 110

Gln Lys Leu Ala Asp Ser Thr Phe Ala Val Asn Gln Phe Asn Leu Leu
115 120 125

Val Ser Asp Gly Ile Ser Val Arg Arg Ser Leu Pro Glu Ile Arg Lys
130 135 140

Pro Ser Cys Arg Asn Met Thr Tyr Pro Asp Asn Leu Pro Thr Thr Ser
145 150 155 160

Val Ile Ile Val Tyr His Asn Glu Ala Tyr Ser Thr Leu Leu Arg Thr
165 170 175

Val Trp Ser Val Ile Asp Arg Ser Pro Lys Glu Leu Leu Lys Glu Ile
180 185 190

Ile Leu Val Asp Asp Phe Ser Asp Arg Glu Phe Leu Arg Tyr Pro Thr
195 200 205

Leu Asp Thr Thr Leu Lys Pro Leu Pro Thr Asp Ile Lys Ile Ile Arg
210 215 220

Ser Lys Glu Arg Val Gly Leu Ile Arg Ala Arg Met Met Gly Ala Gln
225 230 235 240

Glu Ala Gln Gly Asp Val Leu Thr Phe Leu Asp Ser His Cys Glu Cys
 245 250 255
 Thr Lys Gly Trp Leu Glu Pro Leu Leu Thr Arg Ile Lys Leu Asn Arg
 260 265 270
 Lys Ala Val Pro Cys Pro Val Ile Asp Ile Ile Asn Asp Asn Thr Phe
 275 280 285
 Gln Tyr Gln Lys Gly Ile Glu Met Phe Arg Gly Gly Phe Asn Trp Asn
 290 295 300
 Leu Gln Phe Arg Trp Tyr Gly Met Pro Thr Ala Met Ala Lys Gln His
 305 310 315 320
 Leu Leu Asp Pro Thr Gly Pro Ile Glu Ser Pro Thr Met Ala Gly Gly
 325 330 335
 Leu Phe Ser Ile Asn Arg Asn Tyr Phe Glu Glu Leu Gly Glu Tyr Asp
 340 345 350
 Pro Gly Met Asp Ile Trp Gly Gly Glu Asn Leu Glu Met Ser Phe Arg
 355 360 365
 Ile Trp Gln Cys Gly Gly Arg Val Glu Ile Leu Pro Cys Ser His Val
 370 375 380
 Gly His Val Phe Arg Lys Ser Ser Pro His Asp Phe Pro Gly Lys Ser
 385 390 395 400
 Ser Gly Lys Val Leu Asn Thr Asn Leu Leu Arg Val Ala Glu Val Trp
 405 410 415
 Met Asp Asp Trp Lys His Tyr Phe Tyr Lys Ile Ala Pro Gln Ala His
 420 425 430
 Arg Met Arg Ser Ser Ile Asp Val Ser Glu Arg Val Glu Leu Arg Lys
 435 440 445
 Lys Leu Asn Cys Lys Ser Phe Lys Trp Tyr Leu Gln Asn Val Phe Gln
 450 455 460
 Asp His Phe Leu Pro Thr Pro Leu Asp Arg Phe Gly Arg Met Thr Ser
 465 470 475 480
 Ser Ser Asn Ser Ser Val Cys Leu Ala Trp Thr Leu Arg Ser Ser Gly
 485 490 495

Ile Lys Thr Ala Ser Thr Ala Asp Cys Leu Lys Ile Phe His Lys Thr
500 505 510

Gln Leu Trp Leu Tyr Thr Gly Asp Arg Arg Ile Arg Thr Asp Glu His
515 520 525

Leu Cys Leu Ser Val Val Gln Leu Leu His Thr Thr Ser Asp Trp Lys
530 535 540

Ile Gln Leu Lys Glu Cys Ala Gly Phe Asp Thr Glu Tyr Trp Asp Phe
545 550 555 560

Lys Pro Lys Ile Gly Arg Phe Gln Asn Arg Lys Thr Gly Leu Cys Leu
565 570 575

Ala Ser Pro Asp Ile Phe Asp Pro Thr Lys Asp Glu Phe Asn Pro Pro
580 585 590

Ile Val Gln Lys Cys Arg Ser Ser Asn Asp Arg Gln Asn Trp Thr Ile
595 600 605

Thr Glu Met Ser Trp Leu Pro Glu His Pro
610 615

<210> 29
<211> 579
<212> PRT
<213> Caenorhabditis elegans

<400> 29

Met Leu Arg Tyr Ile Ile Pro Arg Lys Lys Gly Thr Phe Val Ile Ala
1 5 10 15

Ala Phe Leu Thr Val Ala Phe Phe Cys Ile Val Ala Tyr His Arg Asn
20 25 30

Asp Arg Arg Arg Thr Lys Phe Gln Phe Pro Asp Ile Glu Lys Tyr Ala
35 40 45

Glu Glu Leu Val Arg Leu Pro Glu Thr Trp Asn Gly Glu Leu His Gln
50 55 60

Ile Pro Asn Tyr Thr Ala Pro Arg Glu Gly Pro Gly Glu Lys Gly Lys
65 70 75 80

Pro Val Val Leu Thr Gly Lys Asp Ala Glu Leu Gly Gln Ala Asp Met

85

90

95

Lys Lys Trp Phe Met Asn Val His Ala Ser Asp Lys Ile Ser Leu Asp
 100 105 110

Arg Asp Val Pro Asp Pro Arg Ile Gln Ala Cys Lys Asp Ile Lys Tyr
 115 120 125

Asp Tyr Ala Ala Leu Pro Lys Thr Ser Val Ile Ile Ile Phe Thr Asp
 130 135 140

Glu Ala Trp Thr Pro Leu Leu Arg Thr Val His Ser Val Ile Asn Arg
 145 150 155 160

Ser Pro Pro Glu Leu Leu Gln Glu Val Ile Leu Leu Asp Asp Asn Ser
 165 170 175

Lys Arg Gln Glu Leu Gln Glu Pro Leu Asp Glu His Ile Lys Arg Phe
 180 185 190

Gly Gly Lys Val Arg Leu Ile Arg Lys His Asp Arg His Gly Leu Ile
 195 200 205

Arg Ala Lys Leu Ala Gly Ala Arg Glu Ala Val Gly Asp Ile Ile Val
 210 215 220

Phe Leu Asp Ser His Cys Glu Ala Asn His Gly Trp Leu Glu Pro Ile
 225 230 235 240

Val Gln Arg Ile Ser Asp Glu Arg Thr Ala Ile Val Cys Pro Met Ile
 245 250 255

Asp Ser Ile Ser Asp Asn Thr Leu Ala Tyr His Gly Asp Trp Ser Leu
 260 265 270

Ser Thr Gly Gly Phe Ser Trp Ala Leu His Phe Thr Trp Glu Gly Leu
 275 280 285

Ser Glu Glu Glu Gln Lys Arg Arg Thr Lys Pro Thr Asp Tyr Ile Arg
 290 295 300

Ser Pro Thr Met Ala Gly Gly Leu Leu Ala Ala Asn Arg Glu Tyr Phe
 305 310 315 320

Phe Glu Val Gly Gly Tyr Asp Glu Glu Met Asp Ile Trp Gly Gly Glu
 325 330 335

Asn Leu Glu Ile Ser Phe Arg Ala Trp Met Cys Gly Gly Ser Ile Glu

